CATGAAGGTTCCTCCTGTCCTGCTTCTCTTTCTTCTGTCCTCAGTGCGAGCTACTGAGCAACCGCAGGTCG ACTTTCTCTGACTGGCAGAACTTCGTGGGCAGGAGACGTTATGGGGCCGAGTCCCAGAACCCCACGGTGAA AGCACTGCTCATCGTGGCCTACTCATTCACCATCGTCTTCTCGCTCTTCGGTAATGTCCTGGTCTGTCATG TCATCTTCAAGAACCAGCGCATGCACTCGGCCACCAGCCTCTTCATTGTCAACCTGGCAGTGGCGGACATC ATGATCACATTGCTCAACACGCCCTTCACTTTGGTCCGCTTTGTGAACAGCACATGGGTGTTTGGGAAGGG CATGTGTCATGTCAGTCGCTTTGCTCAGTACTGTTCTCTACATGTCTCAGCACTGACTCTGACAGCTATCG CAGTGGACCGCCACCAGGTCATCATGCATCCACTGAAGCCTCGGATCTCCATCACCAAGGGTGTCATATAT CAAGTACAGTGAGGACATTGTGCGCTCCCTCTGCCTGCCGGACTTCCCGGAGCCAGCTGACCTCTTCTGGA CGTGTGGCCAAGAAGCTGTGGCTCTGTAACACCATTGGCGACGTGACCACAGAGCAGTACCTCGCCCTGCG ACGCAAGAAGACCACCGTGAAGATGCTGGTGCTTGTGGTAGTCCTCTTTGCCCTCTGCTGGTTCCCTC TCAACTGCTATGTCCTCTTTGTCCAGCAAGGCCATCCACCAACAATGCCCTCTACTTTGCCTTCCAC TGGTTTGCCATGAGCAGTACTTGTTATAACCCCTTCATCTACTGCTGGCTCAATGAGAACTTTAGGGTTGA GCTTAAGGCATTGCTGAGCATGTGCCAAAGGCCACCCAAGCCGCAGGAAGACAGGCTACCCTCCCCAGTTC  $\tt CTTCCTTCAGGGTGGCATGGACAGAGAGAGCCATGGTCGGAGGGCTCCACTACCTAATCACCACTTGCCC$ TCTTCCCAGATCCAGTCTGGGAAGACAGATCTGTCATCTGTGGAACCCGTTGTGGCCATGAGTTAGGGAAA GCTGGAAGTTGGTGGGGGGGGTTCTTTCCTCTCACAATTGACCAGACACTAACAGAGTTGGAAAGTAACA CAGAAGCAGTGAGATGCTTGGGTTCCTAGGAACCTGTCCAGCCCCATCTGATTTGCAAACTTTCTAGAAGA TGCCATGAGGTGTGTGTAGATCTTTGAGCAAGACCTCTGGAAACCACCTCAGCTTCAACAGAGGCTGG TCCAGTCAACCACCTCCAATTGTGTAGCATCTGCCACCTTGCCCTTCCTACTGCTGAGCAACCACAGGGGG ACTTGAGCCATACTATTGGTGGGCCTGCCCCACATGCTCAGAAAAGAACAGGCACAAAGGCTTTCTGAAGT CATTGGAACAGGAATAATCACACAGCTTCAGTGACCTTGGCTCTATCCATGACCAGACAGGACCCATTTTG GCTTCTTAAAAACAAAGAGAAATTAGTATTGCCACTTTGAAAAGTTCAGAAAAGTAAAGAAATGAGTTCAG TAAATTTATCTGTAGCTTTGCCTTCTGTGTGTGTACATTTGTACTTTTAAAATCCTGAACTACACGTGTCC ATGTAGATTGTAATAATTAGCAAGAAACTGGAATATATCAGAGTATTATTGAATTC (SEQ ID NO:1)

MKVPPVLLLFLLSSVRATEQPQVVTEHPSMEAALTGPNASSHFWANYTFSDWQNFVGRRRYGAESQNPTVK ALLIVAYSFTIVFSLFGNVLVCHVIFKNQRMHSATSLFIVNLAVADIMITLLNTPFTLVRFVNSTWVFGKG MCHVSRFAQYCSLHVSALTLTAIAVDRHQVIMHPLKPRISITKGVIYIAVIWVMATFFSLPHAICQKLFTF KYSEDIVRSLCLPDFPEPADLFWKYLDLATFILLYLLPLFIISVAYARVAKKLWLCNTIGDVTTEQYLALR RKKKTTVKMLVLVVVLFALCWFPLNCYVLLLSSKAIHTNNALYFAFHWFAMSSTCYNPFIYCWLNENFRVE LKALLSMCQRPPKPQEDRLPSPVPSFRVAWTEKSHGRRAPLPNHHLPSSQIQSGKTDLSSVEPVVAMS (SEO ID NO:2)

Figure 1

<u>Underlined</u> = deleted in targeting construct

**Bold** = sequence flanking Neo insert in targeting construct

GGGGTGGCAGTCGGCACCATCAGGCTCCCTTGGCGTTTCGGAGTTTTCTCTGTGGTCCCG  ${\tt ACTCTCCGGAGGATCTCGGTTGTCTCCCAAGTCGGAACCTGGCACGGTCCAGGTTCACTC}$ GGAGGTCCGGGCTTCCTCTGTGCCCCGTGCCCCTCGCTCCCAGGCTCCCTCTGTGGTGTG GACTCCTCTAGCCCGGTGCGCTCAGCCCCTCGCACCCAGCCTCCAGGCACAGAGCCCGGC  $\textbf{AGGGAGCTCAGCCCTTGTGCCTAGAGCTGCAGTGGCTGGACATGAAGGTTCCTCCTGT}\underline{\textbf{CC}}$  $\overline{ ext{TGCTTCTTTTCTTGTCCTCAGTGCGAGCTACTGAGCAACCGCAGGTCGTCACTGAGC}}$ ATCCCAGCATGGAGGCAGCCCTGACCGGGCCCAACGCCTCCTCGCACTTCTGGGCCAACT <u>ACACTTTCTCTGACTGGCAGAACTTCGTGGGCAGGAGACGTTATGGGGCCGAGTCCCAGA</u>  ${ t ACCCCACGGTGAAAGCACTGCTCATCGTGGCCTACTCATTCACCATCGTCTTCTCGCTCT}$  ${ t TCGGTAATGTCCTGGTCTGTCATGTCATCTTCAAGAACCAGCGCATGCACTCGGCCACCA}$  ${f GCCTCTTCATTGTCAACCTGGCAGTGGCGGACATCATGATCACATTGCTCAACACGCCCT}$  ${f TCACTTTGGTCCGCTTTGTGAACAGCACATGGGTGTTTGGGAAGGGCATGTGTCA$ <u>GTCGCTTTGCTCAGTACTGTTCTCTACATGTCT</u>CAGCACTGACTCTGACAGCTATCGCAG TGGACCGCCACCAGGTCATCATGCATCCACTGAAGCCTCGGATCTCCATCACCAAGGGTG ACTTCCCGGAGCCAGCTGACCTCTTCTGGAAGTATCTGGACCTGGCCACCTTCATCCTGC TCTACCTACTTCCACTCTTCATTATCTCAGTGGCCTATGCTCGTGTGGCCAAGAAGCTGT GGCTCTGTAACACCATTGGCGACGTGACCACAGAGCAGTACCTCGCCCTGCGACGCAAGA AGAAGACCACCGTGAAGATGCTGGTGCTTGTGGTAGTCCTCTTTTGCCCTCTGCTGGTTCC CTCTCAACTGCTATGTCCTCTTGTCCAGCAAGGCCATCCACACCAACAATGCCCTCT ACTTTGCCTTCCACTGGTTTGCCATGAGCAGTACTTGTTATAACCCCTTCATCTACTGCT GGCTCAATGAGAACTTTAGGGTTGAGCTTAAGGCATTGCTGAGCATGTGCCAAAGGCCAC CCAAGCCGCAGGAAGACAGGCTACCCTCCCCAGTTCCTTCAGGGTGGCATGGACAG AGAAGAGCCATGGTCGGAGGGCTCCACTACCTAATCACCACTTGCCCTCTTCCCAGATCC AGTCTGGGAAGACAGATCTGTCATCTGTGGAACCCGTTGTGGCCATGAGTTAGGGAAAGC  $\tt TGGAAGTTGGTGGGGGGGGGTTCTTTCCTCTCACAATTGACCAGACACTAACAGAGTTGG$ AAAGTAACACAGAAGCAGTGAGATGCTTGGGTTCCTAGGAACCTGTCCAGCCCCATCTGA TGGAAACCACCTCAGCTTCAACAGAGGCTGGTCCAGTCAACCACCTCCAATTGTGTAGCA TCTGCCACCTTGCCCTTCCTACTGCTGAGCAACCACAGGGGGACTTGAGCCATACTATTG GTGGGCCTGCCCACATGCTCAGAAAAGAACAGGCACAAAGGCTTTCTGAAGTCATTGGA ACAGGAATAATCACACAGCTTCAGTGACCTTGGCTCTATCCATGACCAGACAGGACCCAT TTTGGCTTCTTAAAAACAAAGAGAAATTAGTATTGCCACTTTGAAAAGTTCAGAAAAGTA ACATTTGTACTTTTAAAATCCTGAACTACACGTGTCCATGTAGATTGTAATAATTAGCAA GAAACTGGAATATATCAGAGTATTATTGAATTC

Figure 2A

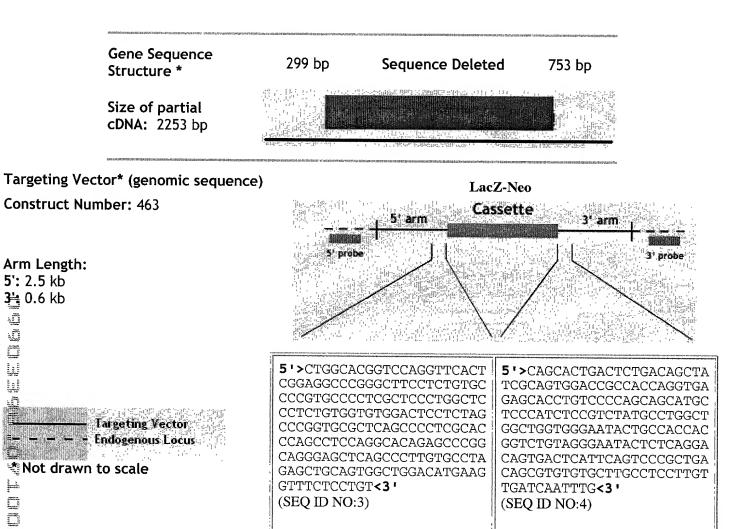


Figure 2B